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AMENDMENT

In the Claims

- 1. (amended) A compound that specifically inhibits the formation of the huburnar C5b-9 complex selected from the group consisting of molecules structurally mimicking CD59 amino acid residues 42 to 58 when they are in a spatial orientation which inhibits formation of the hu C5b 9 complex, wherein the compound is not hu CD59 a peptidomimetic having the structure and function of human CD59 amino acid residues 42-58 of SEO ID NO:3 selected from the group consisting of a peptide, a nucleic acids acid, and a small molecule and an anti-ID anti-CD59 amino acid residues 42-58 antibody, the peptidomimetic and an anti-CD59 antibody binding specifically to human C9 at amino acid residues 26-51 of SEO ID NO:14.
- 2. (amended) The compound of claim 1, selected from the group consisting of proteins, peptides, nucleic acids, and small molecules which bind specifically to amino acids 359 to 384 26-51 of hu human C9 in SEO ID NO:14.
- 3. (amended) The compound of claim 2, wherein the protein compound is an antibody.
- 4. (amended) The compound of claim 2, wherein the protein compound is a chimeric peptide which includes the amino acids 42 to 58 of the human sequence of CD59 in SEO ID NO:3.
- 5. (amended) The compound of claim 2, wherein the peptide compound is a covalently cyclized peptide comprising hu human CD59 amino acid residues 42 to 58 in SEO ID NO:3.

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- (amended) The compound of claim 2, wherein the composition compound 6. is a peptide of less than forty armino acids residues including amino acid residues 42 to 58 of hu human CD59 in SEQ ID NO:3
- (amended) The compound of claim-1, further A composition comprising a 7. compound that specifically inhibits the formation of the human C5b-9 complex selected from the group consisting of a peptidomimetic having the structure and function of human CD59 amino acid residues 42-58 of SEQ ID NO:3 selected from the group consisting of a peptide, a nucleic acids acid, and a small molecule and an anti-ID anti-CD59 amino acid residues 42-58 antibody, the peptidomimetic and an anti-CD59 antibody binding specifically to amino acid residues 26 to 51 of human C9 in SEQ ID NO:14, and a pharmaceutically acceptable carrier for administration to patients in need thercof.
- The compound of claim 1 wherein the compound is a peptidomimetic 8. compound comprising the side chains of hu human CD59 amino acid residues His44, Asn⁴⁸, Asp⁴⁹, Thr⁵¹, Thr⁵², Arg⁵⁵, and Glu⁵⁸ of SEQ ID NO:3 in an equivalent spatial orientation and alignment to that presented on the surface of hu human CD59.
- The compound of claim 8 wherein the spatial orientation and alignment of 9. the side chains of His⁴⁴, Asn⁴⁸, Asp⁴⁹, Thr⁵¹, Thr⁵², Arg⁵⁵, and Glu⁵⁸ of SEQ ID NO:3 in the compound are equivalent to the spatial orientation and alignment deduced by NMR structure determination.
- (three times amended) A method for inhibiting human C5b-9 complex assembly comprising administering to a patient in need thereof an effective amount of a composition comprising a compound binding specifically to amino acid residues 26 to 51

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of human C9 in SEO ID NO:14 selected from the group consisting of a peptidomimetic having the structure and function of human CD59 amino acid residues 42-58 in SEO ID NO:3 selected from the group consisting of proteins, peptides a peptide, a nucleic acids acid, and a small molecule molecules having the structure and function of human CD59 amino acid residues 42-58, and an anti-ID anti-CD59 amino acid residues 42-58 antibody, the peptidomimetic and anti-CD59 antibody binding specifically to amino acid residues 359 384 of human C9.

- (three times amended) The method of claim 10, wherein the compound is 11. a peptidomimetic that is a small molecule which binds specifically to amino acids 359 to 384 26 to 51 of human C9 SEQ ID NO:14.
- 12. (twice amended) The method of claim 10, wherein the protein compound is an antibody.
- (three times amended) The method of claim 10, wherein the protein 13. compound is a chimeric peptide which includes the amino acids 42 to 58 of the human sequence of CD59 in SEQ ID NO:3.
- (three times amended) The method of claim 10, wherein the peptide 14. compound is a covalently cyclized peptide comprising human CD59 amino acid residues 42 to 58 in SEQ ID NO:3.
- (three times amended) The method of claim 10, wherein the 15. peptidomimetic compound is a peptide of less than forty amino acids residues including amino acid residues 42 to 58 of human CD59 in SEQ ID NO:3.

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- (original) The method of claim 10, wherein the composition further 16. comprises a pharmaceutically acceptable carrier for administration to patients in need thereof.
- (once amended) The method of claim 10, wherein the composition is 17. administered to a patient is in need of suppression of complement-mediated inflammation.
- (twice amended) The method of claim 10 wherein the compound is a 18. peptidomimetic comprises comprising the side chains of human CD59 amino acid residues His⁴⁴, Asn⁴⁸, Asp⁴⁹, Thr⁵¹, Thr⁵², Arg⁵⁵, and Glu⁵⁸ of SEQ ID NO:3 in the spatial orientation and alignment of hu human CD59.
- (once amended) The method of claim 18 wherein the spatial orientation 19. and alignment of the side chains of His⁴⁴, Asn⁴⁸, Asp⁴⁹, Thr⁵¹, Thr⁵², Arg⁵⁵, and Glu⁵⁸ of SEQ ID NO:3 in the compound are deduced by NMR structure determination.

Please cancel claims 20-35.